

FIG. 1

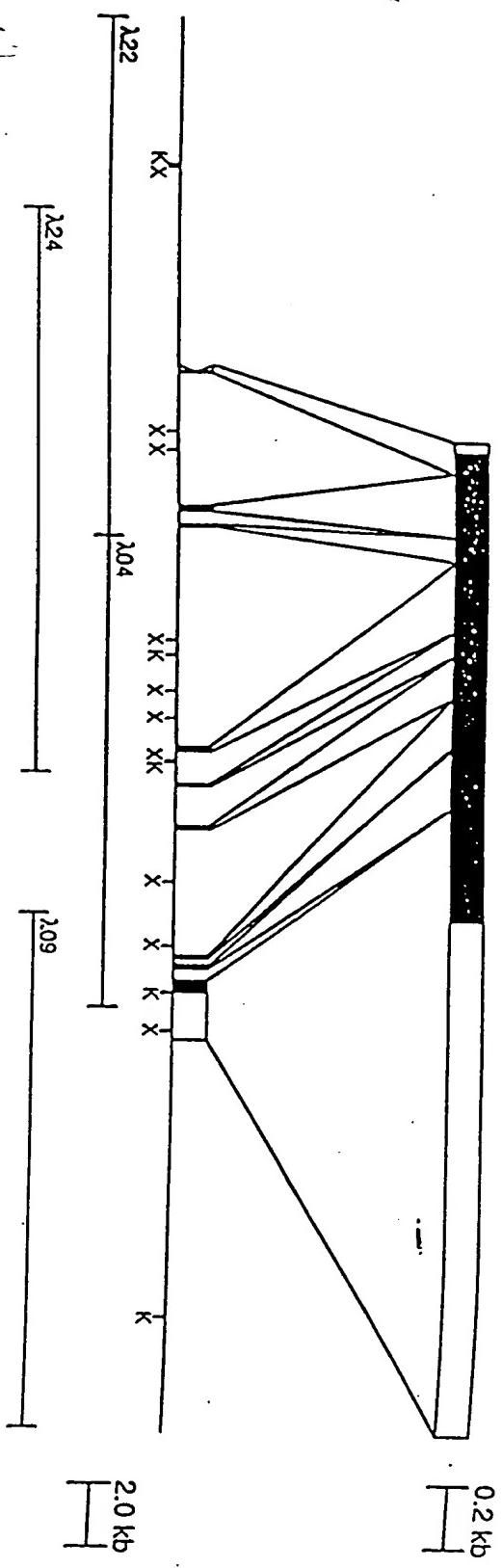


FIG. 2

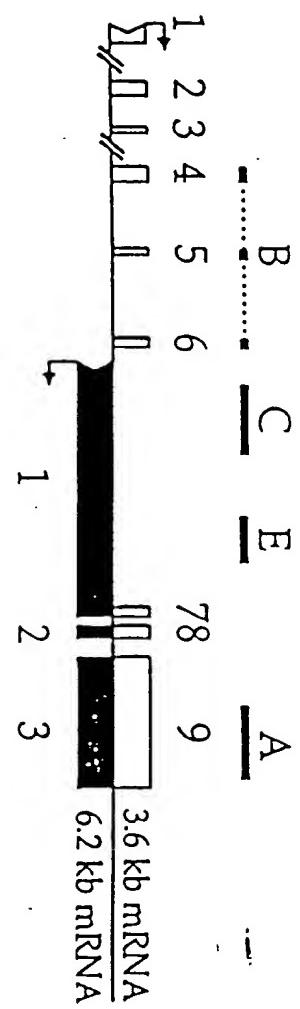


FIG. 3

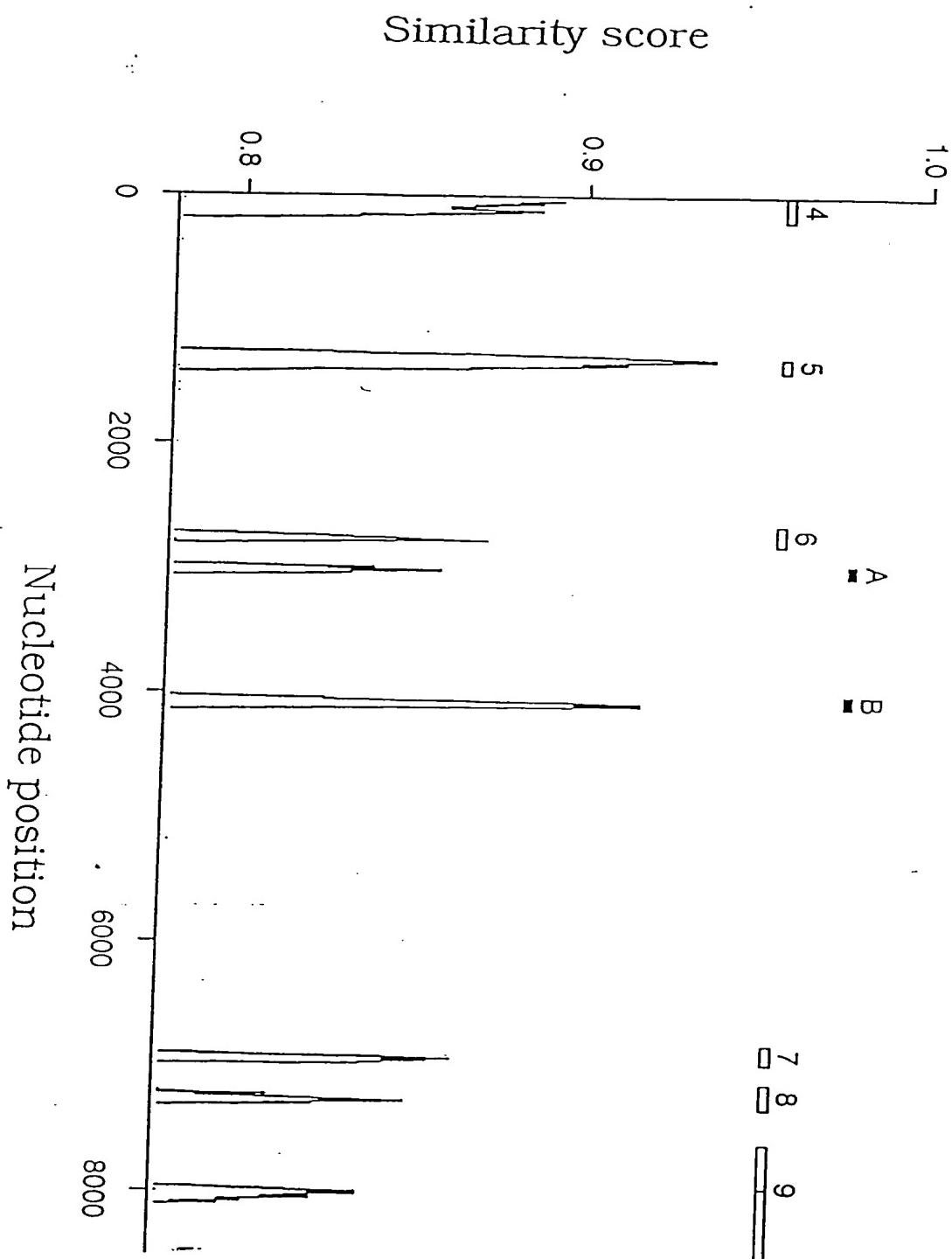


FIG. 4

ε subunit	<u>MSKVVLVPLVGLIGLILQLSRVEGGQOTESKNEASSRDVYVGQPQPOPIENQLSETETGSRVGKULPEASRIL</u>	75
ε subunit	<u>N</u> NTILSNVDHKLRLPGTIGEKPTVVTIELAVNSLGPLSILDNEYTIDLIFSQTWYDERUCYNDTFESLVLNGNVSQ	150
Consensus	L Y RP G S M Y Q W D RL L L	
ε subunit	WIPDTFFFRNSKRTHHEITMPNQMVRKLYKDOCKVLYTIRMTIDAGCSLIMLRFPMDSHSCPLSFSSFSY	225
Consensus	W PDT N H T N R G LY R T C L P D C L SY Y	
ε subunit	KWENFKLEINEKNWKLFQDFDFGVSNKTEELITPVGDFMWTIIFENVSRRFGYVAFQNYPSSVTTMLSWSFN	300
Consensus	W L Q G F L R G Q Y P S VSFN	
ε subunit	IKTESAPARTSLGITSVLMTMIGTSRKNAFPRVSYITALDFYIAICFVFCCFCALLEFAVLNFLLINYQTKAHASP	375
Consensus	AR G TTULMTT R LP A D C FVF AL E	
ε subunit	KLRHPRINSRAHARTRARSRACARQHQEAFCQIVITEGSDGEERPSCSAQQPPSPGSPBGPRLCSKLACCEWC	450
Consensus		
ε subunit	KRFKKVFCMVPDCEGSTWQQGRCLIHVRLDNSRUVFPTFFFENFLYLVCLNL	506
Consensus	D R FP F N YW Y	

FIG. 5A

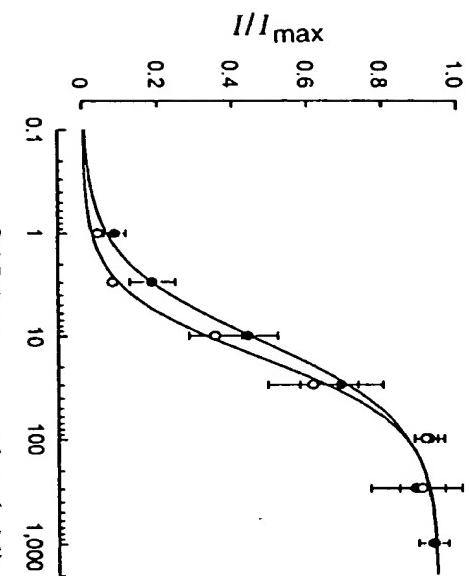


FIG. 5B

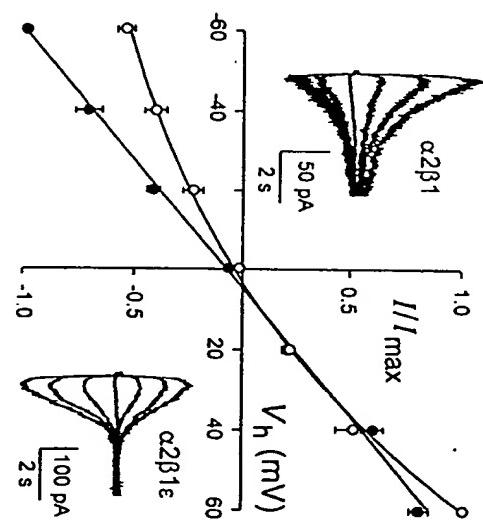


FIG. 5C

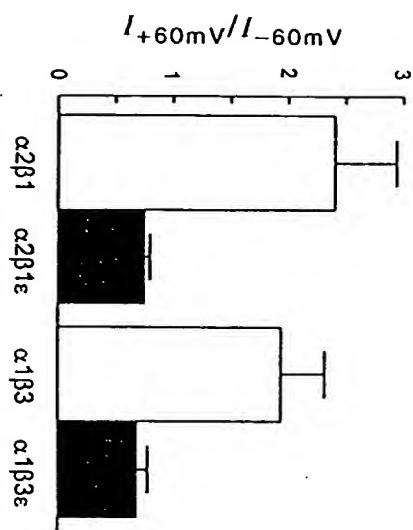


FIG. 6A

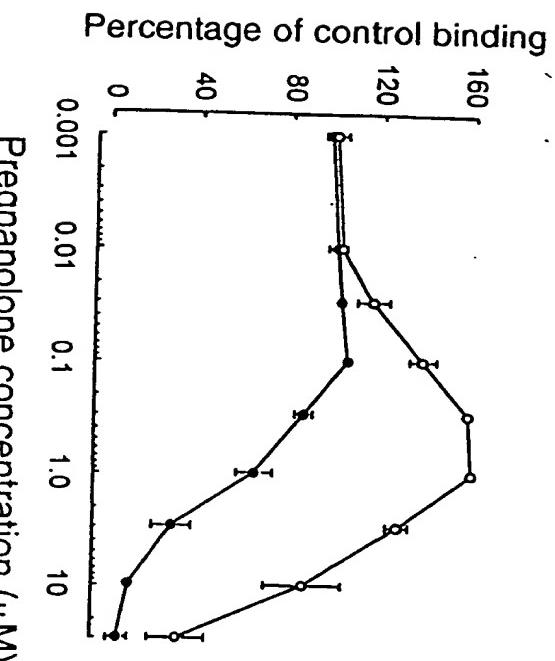


FIG. 6C

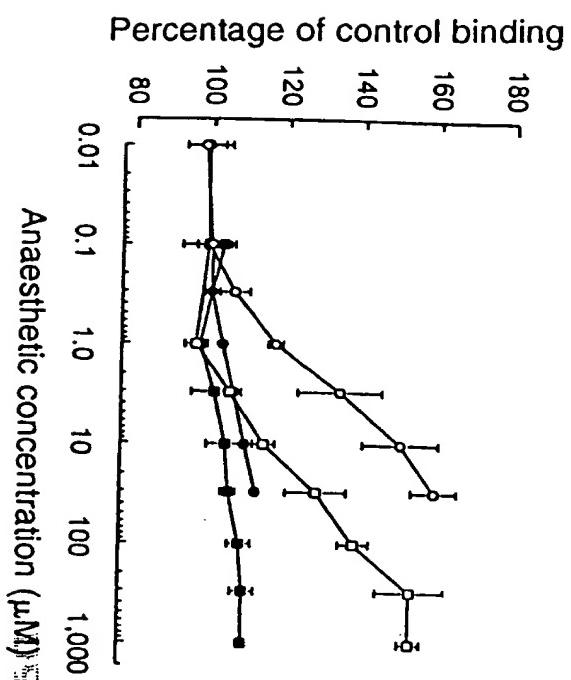


FIG. 6B

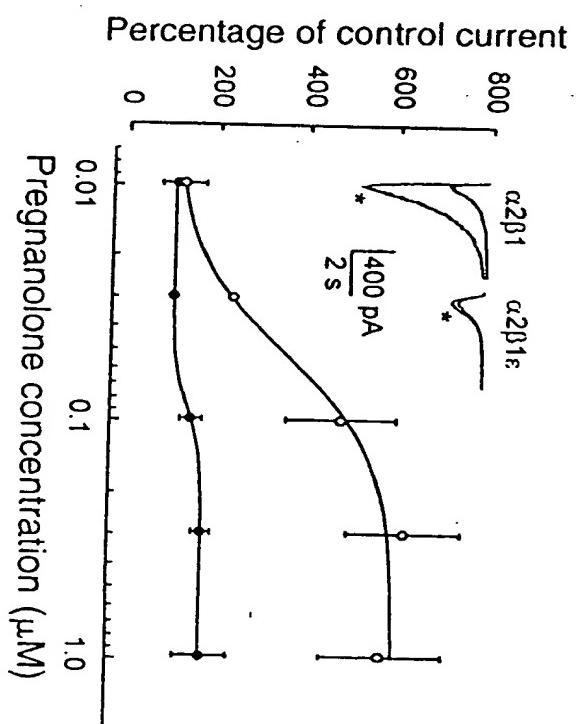
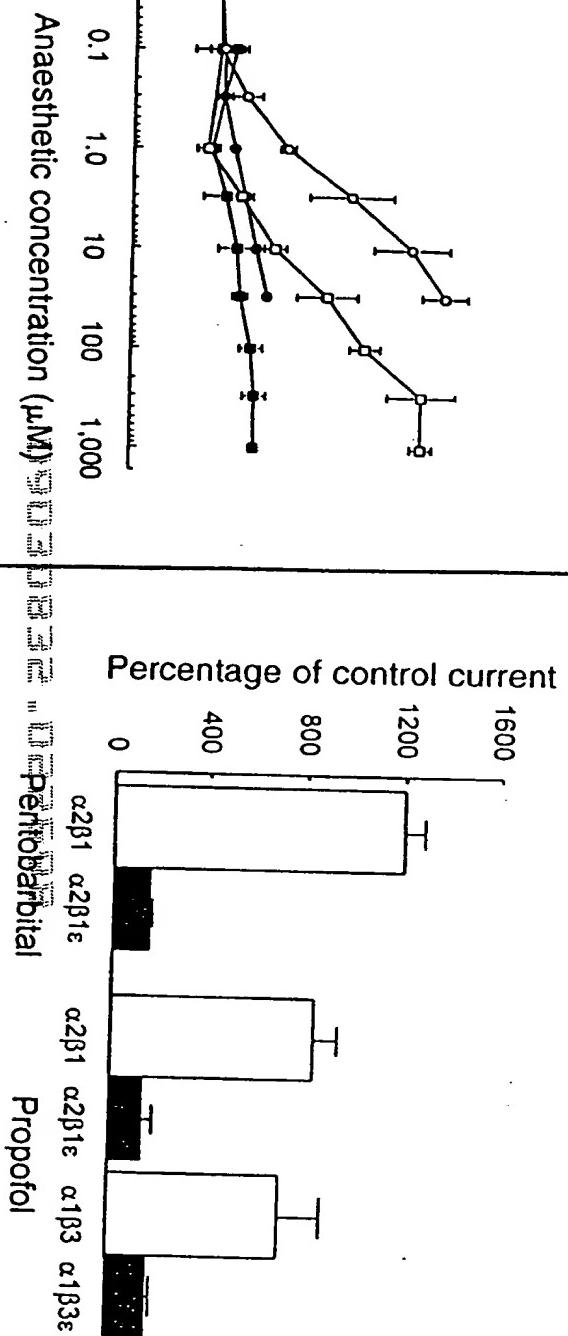
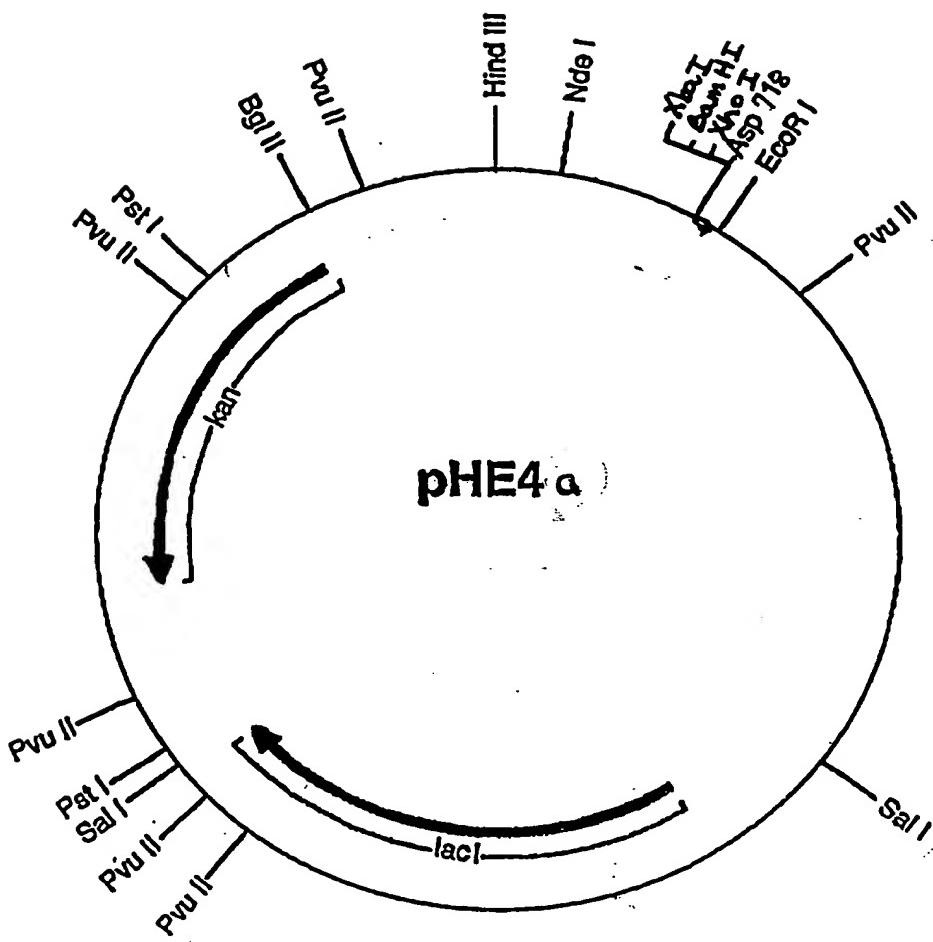


FIG. 6D





**FIG. 7**

FIG. 8

1 AAGCTT AAAA ACTG CAAA ATAG T TTGACTTGATGCACTTAACGAA

-35 Operator 1

-10 Operator 2

50 TAAGATG TACCC ACGCGTGAAGGAATT CACACATTAA

S/D

94 A GAGGAGAAATT A CATATG